

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

GeneCommittee

RRID:SCR_004168

Type: Tool

Proper Citation

GeneCommittee (RRID:SCR_004168)

Resource Information

URL: <http://sing.ei.uvigo.es/GC/>

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Description: Tool for extensively testing the discriminatory power of biologically relevant gene sets in microarray data classification. While the user can work with different gene set collections and several microarray data files to configure specific classification experiments, the tool is able to run several tests in parallel. It is able to render valuable information for diagnostic analyses and clinical management decisions based on systematically evaluating custom hypothesis over different data sets using complementary classifiers, a key aspect in clinical research.

Abbreviations: geneCommittee

Resource Type: production service resource, software resource, data analysis service, analysis service resource, service resource

Defining Citation: [PMID:24475928](https://pubmed.ncbi.nlm.nih.gov/24475928/)

Keywords: dna, microarray, rna-seq, gene expression, classification, gene set, gene, gene set enrichment, functional annotation

Availability: Acknowledgement requested

Resource Name: GeneCommittee

Resource ID: SCR_004168

Alternate IDs: OMICS_02291

Alternate URLs: <https://github.com/michada/GeneCommittee>

Record Creation Time: 20220129T080223+0000

Record Last Update: 20260808T185812+0000

Ratings and Alerts

No rating or validation information has been found for GeneCommittee.

No alerts have been found for GeneCommittee.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.